

Db	241	GGATTGGAAGCGCTCCCTTCTCTCTGTAAGATGACTTTGCATCCCTCCGCTTTGCGAGAG	300
QY	307	TTGGGACAAATTGGATTACTGCTGTGGAAACAAGCCACCACATACTGTAAATPACACTCTT	366
Db	301	TTGGGACAAATTGGATTACTGCTGTGGAAACAAGCCACCACATACTGTAAATPACACTCTT	360
QY	367	CCCTACCAATACCTGACACTGAGGAATGATGAGAGAGGTGCTGCAAAAAGAGCGACACT	426
Db	361	CTTACCAATACCTGACTGAGGAATGATGAGAGAGGTGCTGCAAAAAGAGCGACACT	420
QY	427	CATTCTCTCTACCATCCGCTATCTTCCGACCCTATGAGGCGCATAACTGTGAAACATATG	486
Db	421	CATTCTCTCTACCATCCGCTATCTTCCGACCCTATGAGGCGCATAACTGTGAAACATATG	480
QY	487	GAAGAGCGCTGCTGTATCGGGGCTCTGGAGAACAGACTCGGTATCTACTCTCTCATAC	546
Db	481	GAAGAGCGCTGCTGTATCGGGGCTCTGGAGAACAGACTCGGTATCTACTCTCTCATAC	540
QY	547	AGCCTATGATCTGCGGGCCCAAGGGGCTGACAACAGGTGGGTAAAGGGCTTGGAGCTTG	606
Db	541	AGCCTATGATCTGCGGGCCCAAGGGGCTGACAACAGGTGGGTAAAGGGCTTGGAGCTTG	600
QY	607	TACCTCCAGGCGCCATACATCTCTCCAAAGCTCCCAACTACACCTTACAGAGGGAACACAG	666
Db	601	TACCTCCAGGCGCCATACATCTCTCCAAAGCTCCCAACTACACCTTACAGAGGGAACACAG	660
QY	667	AGTGAATCTCAACGTTAACTACACCCAAAGACTGGACAAAGTCATGTCTGCAGTCAAAAG	726
Db	661	AGTGAATCTCAACGTTAACTACACCCAAAGACTGGACAAAGTCATGTCTGCAGTCAAAAG	720
QY	727	AATGACGSGTCTTCTGTACACTCTCTTTCTTGCTAGAGACTGGTATGAGAAACAACAG	786
Db	721	AATGACGSGTCTTCTGTACACTCTCTTTCTTGCTAGAGACTGGTATGAGAAACAACAG	780
QY	787	GATTAACTGCAATTGTACTCAGAAAGCTTTTGATGCAGGTGGTAGATTTTCTTCCCGGA	846
Db	781	GATTAACTGCAATTGTACTCAGAAAGCTTTTGATGCAGGTGGTAGATTTTCTTCCCGGA	840
QY	847	CAAAACAATTATCGAAGAGAGGAAATCTGTCACTGGAGAAAGCTTTGCTCTCTACATAC	906
Db	841	CAAAACAATTATCGAAGAGAGGAAATCTGTCACTGGAGAAAGCTTTGCTCTCTACATAC	900
QY	907	TGGAATGGAGCGGTTATGCACACTGGATGAATCTGTCTCCGCGCAACCATGATTTGATCG	966
Db	901	TGGAATGGAGCGGTTATGCACACTGGATGAATCTGTCTCCGCGCAACCATGATTTGATCG	960
QY	967	AATAAAAAGAACCTTAAAACTATCTCATATTCGCTTACCGCTGGGGGTGGGGAGAACCTT	1022
Db	961	AATAAAAAGAACCTTAAAACTATCTCATATTCGCTTACCGCTGGGGGTGGGGAGAACCTT	1020
QY	1027	AGAGTCTCAATCAAAAGCTGAGGCCCTGTGCTGTGCTGCTGAGACACAGCTCTCAGAG	1086
Db	1021	AGAGTCTCAATCAAAAGCTGAGGCCCTGTGCTGTGCTGCTGAGACACAGCTCTCAGAG	1080
QY	1087	TGTTGAGGCTACCTTTACCTCACAGGTGAGATGTCCCATCATGATACTTTGGATGTGCG	1144
Db	1081	TGTTGAGGCTACCTTTACCTCACAGGTGAGATGTCCCATCATGATACTTTGGATGTGCG	1144
QY	1147	TTCCCAAGGAATTAATGTCAATCTCTGTGAACACAGCAACACTGAAAGCGGCTTCTTCTT	1206
Db	1141	TTCCCAAGGAATTAATGTCAATCTCTGTGAACACAGCAACACTGAAAGCGGCTTCTTCTT	1200
QY	1207	TGACCTTCGAATATGCTGGATTCTCACTTGAGAAATAAGATAAATATATATCTATCAGA	1266
Db	1201	TGACCTTCGAATATGCTGGATTCTCACTTGAGAAATAAGATAAATATATATCTATCAGA	1260
QY	1267	GACGTGACGGGACCCCTCTTCAGGCTGTAAATTTGAGAAACATCAGGATACACACTTC	1324
Db	1261	GACGTGACGGGACCCCTCTTCAGGCTGTAAATTTGAGAAACATCAGGATACACACTTC	1318

```

Sequence 12938, Application US/10198846
Publication NO. US20030099974A1
GENERAL INFORMATION:
APPLICANT: Lillie, James
APPLICANT: Xu, Yongyao
APPLICANT: Wang, Youzhen
APPLICANT: Steinhmann, Kathleen
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
TITLE OF INVENTION: FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
TITLE OF INVENTION: THERAPY OF BREAST CANCER
FILE REFERENCE: MRI-049
CURRENT APPLICATION NUMBER: US/10/198, 846
CURRENT FILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: 60/306, 220
PRIOR FILING DATE: 2001-07-18
NUMBER OF SEQ ID NOS: 14084
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 12938
LENGTH: 1614
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 1613..1614
OTHER INFORMATION: n = A,T,C or G
US-10-198-846-12938

```

Query Match	Similarity	Score	DB	Length
Match	100.0%	Pred. NO. 1.4e-303		
Matches	574: Conservative	0: Mismatches	0: Indels	0: Gaps
QY	490	GGAGCGCGCTGGTGTATCCGGGCTCTGGAGAAACAGATGGGTATCTACTCTCCCTACAGC	549	
DB	678	GGAGCGCGCTGGTGTATCCGGGCTCTGGAGAAACAGATGGGTATCTACTCTCCCTACAGC	737	
QY	550	CTATGATGCTGCGGCCCAAGGGCGTCACAACTGGTGGCTTAAAGGCTTGGAGCTTGTAC	609	
DB	738	CTATGATGCTGCGGCCCAAGGGCGTCACAACTGGTGGCTTAAAGGCTTGGAGCTTGTAC	797	
QY	610	CTCCAGGCGCATACATCCCTTCCAAAGCTCCCACTACCCCTACAGAGGAAACACAGAGT	669	
DB	798	CTCCAGGCGCATACATCCCTTCCAAAGCTCCCACTACCCCTACAGAGGAAACACAGAGT	857	
QY	670	AGAAATCAACGTTAACTACACCAACAAGACTGGACAAAGTCATGCTCGAGTGAAGGAT	729	
DB	858	AGAAATCAACGTTAACTACACCAACAAGACTGGACAAAGTCATGCTCGAGTGAAGGAT	917	
QY	730	TGAGGGGTTCCTGTCACCTCTTTTTCGTCAGAGACTGGTAAATAGGAACAAACAGGAT	789	
DB	918	TGAGGGGTTCCTGTCACCTCTTTTTCGTCAGAGACTGGTAAATAGGAACAAACAGGAT	977	
QY	790	TAACTGTAATGTACTCGAAGGCTTGTGATGCAGGTGGTAGATTTTCTTCCCGGAACA	849	
DB	978	TAACTGTAATGTACTCGAAGGCTTGTGATGCAGGTGGTAGATTTTCTTCCCGGAACA	1037	
QY	850	ACAACCTTATATAGAAAGACGAAATTCGTCACTGAGAAAGCTTTCCTTCTACTACTGG	909	
DB	1038	ACAACCTTATATAGAAAGACGAAATTCGTCACTGAGAAAGCTTTCCTTCTACTACTGG	1097	
QY	910	AATGGAGCGGTTATGACACACTGGATGAAATCTGTCTCCCTGGCAACCAATGATGTGAT	969	
DB	1098	AATGGAGCGGTTATGACACACTGGATGAAATCTGTCTCCCTGGCAACCAATGATGTGAT	1157	
QY	970	AAAAAGACCTAAAACTATCTCATATTCGTAGCCCTTGGGTGGGAGAACCTTGA	1029	
DB	1158	AAAAAGACCTAAAACTATCTCATATTCGTAGCCCTTGGGTGGGAGAACCTTGA	1217	
QY	1030	GTCCTCAAGTCAAAAGTCGCGCCGCTGGCGGT	1063	
DB	1218	GTCCTCAAGTCAAAAGTCGCGCCGCTGGCGGT	1251	

RESULT 2
US-10-198-846-12938

RESULT 3

Query Match	13.4%	Score 208;	DB 9;	Length 208;
Best Local Similarity	100.0%	Pred. No. 4.7e-103;		
Matches 208; Conservative	0;	Mismatches	0;	Indels 0; Gaps 0

[illegible]

```

RESULT 7
US-10-198-846-9256
; Sequence 9256, Application US/10198846
; Publication NO. US2003009974A1
GENERAL INFORMATION:
APPLICANT: Lillie, James
APPLICANT: Xu, Yongyao
APPLICANT: Wu, Youzhen
APPLICANT: Steilmann, Kathleen
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
TITLE OF INVENTION: FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
FILE OF INVENTION: MRI-049
FILE REFERENCE: MRI-049
CURRENT APPLICATION NUMBER: US/10/198,846
CURRENT FILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: 60/306,220
PRIOR FILING DATE: 2001-07-18
NUMBER OF SEQ ID NOS: 14084
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 9256
LENGTH: 638
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 2, 4, 16, 525, 528, 557, 566, 576, 607, 614
OTHER INFORMATION: n = A,T,C or G
US-10-198-846-9256

```

Query Match	12.1%;	Score 188;	DB 14;	Length 638;
Best Local Similarity	100.0%;	Pred. No. 4.3e-92;		
Matches 188;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0

QY	1	GTGATTGTTATCTTGGTGTGCGAGGAGCAGCAGAGAAAGGAGATTGGSTCGAANAATCGC	60
Db	80	GTATATTGTTATCTTGGTGTGCGAGGAGCAGCAGAGAAAGGAGATTGGSTCGAANAATCGC	135
QY	61	CCTGCCGCACACAGACACAGCCCACTAGTGGGACAGGGGTCCTGACTAGACTTAACGTGG	120
Db	140	CTGCGCCGACACAGACACAGCCCACTAGTGGGACAGGGGTCCTGACTAGACTTAACGTGG	195
QY	121	CTGTGTCCTGCTGTTTTTCACTGTCCCTGGAAAAGCCCTGAATGGCACTGAAATGAGCA	180
Db	200	CTGTGTCCTGCTGTTTTTCACTGTCCCTGGAAAAGCCCTGAATGGCACTGAAATGAGCA	255
QY	181	TAGATGAG	188
Db	260	TAGATGAG	267

RESULT 8

```

US-10-027-632-161386/c
; Sequence 161386, Application US/10027632
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.

```

Query Match 3.28; Score 50; DB 13; Length 817;
Best Local Similarity 100.0%; Pred. No. 1.7e-16;
Matches 50; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1035 AAGTCAAGTCGTGGCCCTGTGTGCTGTTCTGGAGCAGCGTTCTGCAG 1084
|||||
Db 71 AAGTCAAGTCGTGGCCCTGTGTGCTGTTCTGGAGCAGCGTTCTGCAG 22

```

RESULT 9
US-10-027-632-161387/c
; Sequence 161387, Application US/10027632
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
; TITLE OF INVENTION: Polymorphisms in the Human Genome
; FILE REFERENCE: 108827.129
; CURRENT APPLICATION NUMBER: US/10/027,632
; CURRENT FILING DATE: 2002-04-30
; PRIOR APPLICATION NUMBER: US 60/218,006
; PRIOR FILING DATE: 2000-07-12
; PRIOR APPLICATION NUMBER: US 60/198,676
; PRIOR FILING DATE: 2000-04-20
; PRIOR APPLICATION NUMBER: US 60/193,483
; PRIOR FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: US 60/185,218
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/167,363
; PRIOR FILING DATE: 1999-11-23
; PRIOR APPLICATION NUMBER: US 60/156,358
; PRIOR FILING DATE: 1999-09-28
; PRIOR APPLICATION NUMBER: US 60/146,002
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 325720
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 161387
; LENGTH: 817
; TYPE: DNA
; ORGANISM: Human
US-10-027-632-161387

```

Query Match	3.28; Score 50; DB 13; Length 817;
-------------	------------------------------------

Best Local Similarity 100.0%, Pred. No. 1.7e-16;
Matches 50; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
DB 1035 AAGTAAAGTCGTGGCCCTGTGTGCTGGTTCGGAGAGCGTTCGAG 1084
11 AAGTAAAGTCGTGGCCCTGTGTGCTGGTTCGGAGAGCGTTCGAG 22

RESULT 10
US-09-783-590-8477
Sequence 8477, Application US/09783590
Patent No. US20020110850A1
GENERAL INFORMATION:
APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, William A.
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE REFERENCE: PO-16.2C1
CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 8477
LENGTH: 501
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (10)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (48)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (98)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (141)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (173)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (229)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (315)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (358)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (376)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (391)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (404)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (411)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (427)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature

LOCATION: (430)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (461)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (465)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (467)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (473)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (479)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (487)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (492)
OTHER INFORMATION: n equals a,t,g, or c
US-09-783-590-8477

Query Match
Best Local Similarity 3.2%; Score 49; DB 10; Length 501;
Matches 49; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 140 ACTGTCCTGGAAGGCGCTGGAAGTGCGACTGGAATGAGCATATGATGAG 168
DB 230 ACTGTCCTGGAAGGCGCTGGAAGTGCGACTGGAATGAGCATATGATGAG 278

RESULT 11
US-10-177-900-10/c
Sequence 10, Application US/10177900
Publication No. US20030068787A1
GENERAL INFORMATION:
APPLICANT: Jackson, Jennifer L.
APPLICANT: Corley, Neil C.
APPLICANT: Guegler, Karl J.
APPLICANT: Arvizu, Chandra
TITLE OF INVENTION: AN ANTIBODY SPECIFICALLY BINDING CYCLOPHILIN-TYPE PEPTIDYL-PRO
FILE REFERENCE: PF-0582-2 CIP
CURRENT APPLICATION NUMBER: US/10/177,900
CURRENT FILING DATE: 2002-06-20
PRIOR APPLICATION NUMBER: 09/440,828
PRIOR FILING DATE: 1999-11-15
PRIOR APPLICATION NUMBER: 09/136,442
PRIOR FILING DATE: 1998-08-19
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PERL Program
SEQ ID NO 10
LENGTH: 2922
TYPE: DNA
ORGANISM: Mus musculus
FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No. US20030068787A1 023075_Mm.3
US-10-177-900-10

Query Match
Best Local Similarity 2.4%; Score 38; DB 14; Length 2922;
Matches 38; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 347 CATACTGTAAATACACTCTTCCTGACCAATGACCTGAC 384
DB 1149 CATACTGTAAATACACTCTTCCTGACCAATGACCTGAC 1112

RESULT 12

US-09-960-352-10848
; Sequence 10848, Application US/09960352
; Patent No. US20020137139A1
; GENERAL INFORMATION:
; APPLICANT: Warren, Wesley C.
; APPLICANT: Rao, Mengding
; APPLICANT: Byatt, John C.
; APPLICANT: Mathalagan, Nagappan
; TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
; FILE REFERENCE: 16511.006/37-21(10298)C
; CURRENT APPLICATION NUMBER: US/09/960,352
; CURRENT FILING DATE: 2001-09-24
; NUMBER OF SEQ ID NOS: 15112
; SEQ ID NO 10848
; LENGTH: 394
; TYPE: DNA
; ORGANISM: Bos taurus
; OTHER INFORMATION: Clone ID: 46-LIB34-084-Q1-E1-D6
US-09-960-352-10848

Query Match 1.9%: Score 29; DB 10; Length 394;
Best Local Similarity 100.0%; Pred. No. 5.5e-05;
Matches 29; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 308 TGGGACATGTTGGATTCTGTGGAC 336
|||||
DB 182 TGGGACATGTTGGATTCTGTGGAC 210

RESULT 13
; Sequence 162668, Application US/10027632
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
; FILE REFERENCE: 108827.129
; CURRENT APPLICATION NUMBER: US/10/027,632
; CURRENT FILING DATE: 2002-04-30
; PRIOR APPLICATION NUMBER: US 60/218,006
; PRIOR FILING DATE: 2000-07-12
; PRIOR APPLICATION NUMBER: US 60/198,676
; PRIOR FILING DATE: 2000-04-20
; PRIOR APPLICATION NUMBER: US 60/193,483
; PRIOR FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: US 60/185,218
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/167,363
; PRIOR FILING DATE: 1999-11-23
; PRIOR APPLICATION NUMBER: US 60/156,358
; PRIOR FILING DATE: 1999-09-28
; PRIOR APPLICATION NUMBER: US 60/146,002
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 325720
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 162668
; LENGTH: 765
; TYPE: DNA
; ORGANISM: Human
US-10-027-632-162668

Query Match 1.7%: Score 26; DB 13; Length 765;
Best Local Similarity 100.0%; Pred. No. 0.0025;
Matches 26; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 269 TCCTGATGACTTTCATCCCTCTC 294
|||||
DB 345 TCCTGATGACTTTCATCCCTCTC 370

RESULT 14
US-10-311-455-568/C

; Sequence 568, Application US/10311455
; Publication No. US20030143606A1
; GENERAL INFORMATION:
; APPLICANT: Olex, Alexander
; APPLICANT: PIEPENBROCK, Christian
; APPLICANT: BERLIN, Kurt
; TITLE OF INVENTION: Diagnosis of Diseases Associated with the Immune System by Det
; FILE REFERENCE: 5013.1014
; CURRENT APPLICATION NUMBER: US/10/311,455
; CURRENT FILING DATE: 2002-12-16
; PRIOR APPLICATION NUMBER: PCT/EP01/07537
; PRIOR FILING DATE: 2001-07-02
; PRIOR APPLICATION NUMBER: DE 10032529.7
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: DE 10043826.1
; PRIOR FILING DATE: 2000-09-01
; NUMBER OF SEQ ID NOS: 2424
; SEQ ID NO 568
; LENGTH: 11976
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-10-311-455-568

Query Match 1.4%: Score 22; DB 12; Length 11976;
Best Local Similarity 100.0%; Pred. No. 0.4;
Matches 22; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1473 ATATACACCATTAATAA 1494
|||||
DB 1472 ATATACACCATTAATAA 1451

RESULT 15
; Sequence 29392, Application US/10027632
; GENERAL INFORMATION:
; APPLICANT: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
; FILE REFERENCE: 108827.129
; CURRENT APPLICATION NUMBER: US/10/027,632
; CURRENT FILING DATE: 2002-04-30
; PRIOR APPLICATION NUMBER: US 60/218,006
; PRIOR FILING DATE: 2000-07-12
; PRIOR APPLICATION NUMBER: US 60/198,676
; PRIOR FILING DATE: 2000-04-20
; PRIOR APPLICATION NUMBER: US 60/193,483
; PRIOR FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: US 60/185,218
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/167,363
; PRIOR FILING DATE: 1999-11-23
; PRIOR APPLICATION NUMBER: US 60/156,358
; PRIOR FILING DATE: 1999-09-28
; PRIOR APPLICATION NUMBER: US 60/146,002
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 325720
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 29392
; LENGTH: 586
; TYPE: DNA
; ORGANISM: Human
US-10-027-632-29392

Query Match 1.3%: Score 20; DB 13; Length 586;
Best Local Similarity 100.0%; Pred. No. 4.8;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 23 AGAGCAGCAGAGAGAGAG 42
|||||

Db 149 AGAGGACAGCAGAGAGGAG 130

Search completed: August 23, 2003, 18:28:30
Job time : 283 secs
